

Multitrophic interaction facilitates parasite–host relationship between an invasive beetle and the honey bee

Baldwyn Torto*, Drion G. Boucias†, Richard T. Arbogast‡, James H. Tumlinson§¶, and Peter E. A. Teal†¶

*International Centre of Insect Physiology and Ecology, P.O. Box 30772-00100, Nairobi, Kenya; †Institute of Food and Agricultural Sciences, Department of Entomology and Nematology, University of Florida, 110620, Gainesville, FL 32611; ‡Center for Medical, Agricultural, and Veterinary Entomology, U.S. Department of Agriculture–Agricultural Research Service, 1700 SW 23 Drive, Gainesville, FL 32608; and §Center for Chemical Ecology, Department of Entomology, Pennsylvania State University, University Park, PA 16802

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Colony defense by honey bees, *Apis mellifera*, is associated with stinging and mass attack, fueled by the release of alarm pheromones. Thus, alarm pheromones are critically important to survival of honey bee colonies. Here we report that in the parasitic relationship between the European honey bee and the small hive beetle, *Aethina tumida*, the honey bee's alarm pheromones serve a negative function because they are potent attractants for the beetle. Furthermore, we discovered that the beetles from both Africa and the United States vector a strain of *Kodamaea ohmeri* yeast, which produces these same honey bee alarm pheromones when grown on pollen in hives. The beetle is not a pest of African honey bees because African bees have evolved effective methods to mitigate beetle infestation. However, European honey bees, faced with disease and pest management stresses different from those experienced by African bees, are unable to effectively inhibit beetle infestation. Therefore, the environment of the European honey bee colony provides optimal conditions to promote the unique bee–beetle–yeast–pollen multitrophic interaction that facilitates effective infestation of hives at the expense of the European honey bee.

alarm pheromone | kairomone | small hive beetle

Interactions between species play central roles in evolution, and most species can be defined by interspecific interactions (1). For example, the ability of parasites and predators to survive and reproduce depends on the ability to overcome the host defenses, thereby increasing their fitness advantage (1, 2). Conversely, hosts are constantly evolving ways to defend against attack. Normally a tight balance exists in these interspecies interactions, allowing both host and attacker to survive. Only when the attacking species is freed from the constraints of host defenses does the balance shift in favor of the attacker. An example of such interactive coevolution occurs between the African honey bee (AHB) and the small hive beetle (SHB), a facultative parasite. In the native range, subSaharan Africa, the SHB is a minor pest of bee hives (3–5) because the AHB has evolved effective behavioral ways to control infestation including removal of eggs of the beetle from comb cells as a form of hygienic behavior and imprisonment and encapsulation of the invading beetles by guard bees into cracks and crevices in the hive (6). The beetles, in turn, survive imprisonment by behavioral mimicry involving tactile stimuli to initiate their feeding by trophallaxis from the guard bees (6).

The SHB was recently introduced into the United States and Australia and has become a devastating pest of resident European honey bees (EHB) (5, 7, 8), and consequently, is a threat to EHB pollinated crops, worth \$14 billion per annum in the United States. Given the similar behavioral imprisonment response by EHBs to invading beetles (6), the biological interactions between the EHB and the SHB that contribute to the beetle's highly invasive parasitic relationship are unclear. Adult

beetles are attracted to volatiles of EHBs (9, 10), and we know that the attraction is mediated by a blend of components dominated by the honey bee's alarm pheromones (10), including isopentyl acetate (IPA), 2-heptanone, and methyl benzoate, which account for ≈70–80% of the blend. Interestingly, the sting response and alarm pheromone are the key components of the honey bee defense system against predators and parasites (11). Fitness advantages for the bees in releasing alarm pheromones include triggering mass attack against an intruder, recruitment of more guard bees, and possibly sending a signal to repel would-be intruders in the vicinity (11). Little is known regarding the risk involved in the release of alarm pheromones. However, the critical importance of alarm pheromones to honey bee survival, coupled with the fact that the SHB is attracted to a blend of chemicals dominated by the honey bee alarm pheromones, suggests that they could provide a unique cue for SHB attack (1, 2, 11). Using bioassays plus chemical and molecular analytical techniques, we found a unique semiochemically mediated multitrophic interaction based on honey bee alarm pheromones. This relationship involves the honey bee, small hive beetle, a yeast species vectored by the beetle, and bee-collected pollen, resulting in a significant threat to survival of the EHB, already faced with multiple management stresses, after beetle invasion.

Results and Discussion

Parasitic Beetle Detects Alarm Pheromone Released at Entrance of Unstressed Honey Bee Colony. GC-MS analysis revealed that 100 EHBs ($n = 3$) stressed artificially, by confinement in a container, released ≈1,500- to 10,000-fold more alarm pheromone as indicated by release of isopentyl acetate than released at the entrance of the undisturbed honey bee colony ($n = 3$), estimated to contain 40,000–60,000 bees (12) (50–120 ng/hr released by the artificially stressed 100 worker bees vs. 0.8–6 ng/hr released by the undisturbed honey bee colony). In coupled gas chromatographic-electroantennogram (GC-EAD) analyses, antennae of either sex of the beetle ($n = 5$ male and 5 female antennae) detected the equivalent of 2 ng of IPA in the volatiles captured at the entrance of the undisturbed honey bee colony ($n = 3$ honey

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Abbreviations: AHB, African honey bee; EHB, European honey bee; GC-EAD, gas chromatographic-electroantennogram; IPA, isopentyl acetate; SDAY, Sabouraud dextrose agar yeast; SHB, small hive beetle.

¶To whom correspondence may be addressed. E-mail: peter.teal@ars.usda.gov. or jht2@psu.edu.

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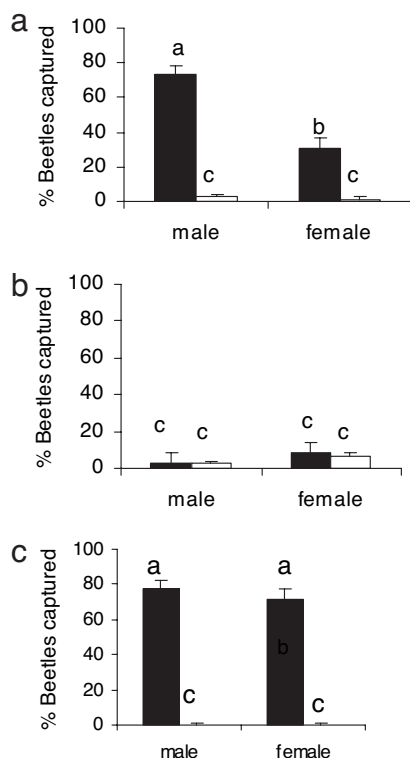


Fig. 4. Wind tunnel responses of *A. tumida* males and females (4–8 weeks old) to volatiles released from different media inoculated with (filled bars) or without (open bars) the *Kodamaea* strain isolate from beetle larvae sterilized bee collected pollen (a), sterilized commercial pollen substitute (Bee Pro) (b); and sterilized bee collected pollen (c) (filled bars) compared with SDAY (open bars), both inoculated with the yeast isolate. There were three replicates in each test. $n = 25$ beetles per replicate. Bars with the same letter were not significantly different ($P < 0.05$, LSD test).

and (ii) brood. We then tested the attractiveness of the volatiles released from the diets that had or had not been fed on by the beetles in flight tunnel assays. Surprisingly, only the pollen diet fed on by either sex of the SHB for three days was highly attractive [supporting information (SI) Table 1], and a major chemical component of these volatiles was IPA along with other alarm pheromones and fermentation-related products.

Because nitidulid beetles are well known vectors of fungi (13), we investigated the possible involvement of a fungus in the production of IPA in the volatiles of the SHB-infested comb. We plated homogenates from both larvae and adults collected in the United States on a commonly used substrate for fungal growth, Sabouraud dextrose agar yeast (SDAY) extract. Microscopic examination of colonies developed on the plates revealed the presence of budding yeast cells. We then grew the yeast on three different sterilized substrates: pollen collected from traps placed at the entrances of honey bee hives (bee-collected pollen); a commercial pollen substitute (Bee Pro, Hackensack, MN), widely used in the beekeeping industry; and SDAY media. Comparison of responses of SHBs to volatiles released from the different sterilized media inoculated with or without the yeast isolate in a wind tunnel, revealed significant differences (Fig. 4; $F_{(3, 11)} = 45.9$, $r^2 = 0.95$, $P = 0.0001$). Significantly more SHBs were lured into traps releasing volatiles from the yeast-inoculated pollen than to control traps (Fig. 4a). In contrast, there were no significant differences in the numbers of beetles lured into traps releasing volatiles from yeast-free commercial pollen substitute and yeast-inoculated commercial pollen substitute (Fig. 4b), and neither did the beetles respond to volatiles

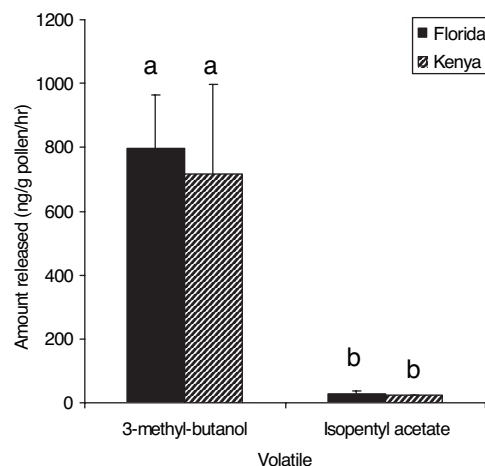


Fig. 5. Amount of 3-methyl-butanol (honey bee alarm pheromone precursor) and IPA (honey bee alarm pheromone) released by sterilized bee-collected pollen inoculated for 7 days with the yeast *K. ohmeri*, isolated from *Aethina tumida* obtained from honey bee colonies in Florida and Kenya (mean of three replicates, error bars represent standard error). Bars with the same letter are not significantly different (Student's *t* test, $P < 0.05$).

released from yeast-free SDAY media and the yeast-inoculated SDAY media. When the beetles were given a choice between volatiles released from yeast-inoculated-bee-collected pollen and those from yeast-inoculated SDAY media, significantly more beetles were captured in the trap releasing volatiles from the yeast-inoculated bee-collected pollen (Fig. 4c). GC-MS analysis of volatiles released from bee-collected pollen inoculated with the yeast revealed that the honey bee alarm pheromone (IPA) was consistently present in the volatiles being released at a rate of $\approx 20 \text{ ng} \cdot \text{g} \text{ of pollen}^{-1} \cdot \text{hr}^{-1}$ (Fig. 5). IPA was absent from the volatiles of other sources. In addition, we confirmed the presence of ethyl esters which are also known alarm pheromone mimics (14) in volatiles from the yeast-inoculated pollen.

It was clear that yeast associated with the U.S. population of SHB produced bee alarm pheromones attractive to the beetles. What was unclear was whether the yeast was resident only in the U.S. population of SHB or it was present in all populations of SHB. We addressed this by growing yeast colonies from beetles collected from colonies of the AHB in Kenya. Microbial isolation coupled with DNA extractions and sequencing showed that both the U.S. and African beetles vectored the same yeast, identified as a strain of *Kodamaea ohmeri*. Additionally, analysis of volatiles released from yeast of both strains produced the same volatiles when incubated with bee-collected pollen (Fig. 5). Therefore, yeast in colonies of AHB colonies would produce the same attractants as are produced in EHB colonies. This indicates that SHBs will attack honey bee colonies, whether strong or weak and irrespective of the subspecies, European or African.

Our findings suggest that a SHB attack on EHB colonies likely proceeds as follows: First, initial SHB infestation of the honey bee colony is caused by the beetle detecting colony volatiles, including alarm pheromones at thresholds lower than detected by worker bees. The beetle associates these chemicals with the presence of food resources in the colony. This initial attack also could be aided inadvertently by the bees themselves when they collect pollen from flowers contaminated by yeast spores deposited by flower-feeding nitidulids. Indeed, yeasts of the genus *Kodamaea* have been found in certain ephemeral flowers, which serve as breeding and feeding sites for nitidulid beetles (15). Unlike in healthy colonies, once initial infestation occurs, the invading beetles escape confinement from guard bees in colonies

SDAY. Inoculated and noninoculated plates were incubated at 28°C for 7 days before use. For GC-EAD analysis, 5- μ l aliquots of the volatile extracts were analyzed (GC-EAD 2000; Syntech, Hilversum, The Netherlands) on a Hewlett-Packard 5890 Series II gas chromatograph equipped with a HP-5 column (30 m \times 0.32 mm ID \times 0.25 μ m) (Agilent), as described in ref. 10. The oven temperature was held at 35°C for 5 min, then programmed to increase at 10°C/min to 220°C and held at this temperature for 5 min. For EAD, excised antennae of either male or female beetles or worker bees were held between gold electrodes in conductivity gel (Syntech).

Isolation of Yeast Strain. Larvae of the SHB, removed from an infested hive of EHBs in Florida, were surface-sterilized (70% ethanol) for \approx 30 seconds and then rinsed twice in sterile water. Larvae from a strain of SHBs were obtained from AHB colonies maintained at the International Center of Insect Physiology and Ecology, and they were similarly treated. Larvae were homogenized in sterile water (1 insect/ml) and streaked for isolation on SDAY. Honey samples also were collected from the infested hive and plated for isolation on SDAY. Inoculated plates were incubated at 31°C for 1–3 days. Individual colonies were selected and subcultured on SDAY. Isolates were inoculated into Durham tubes containing autoclaved bee pollen broth (1% aqueous pollen) tubes and incubated at 31°C for 5 days. The strain NRRL Y27634 (ARS Culture Collection) selected from the gas-producing isolates produced a colony morphology characteristic of the majority of yeast colonies observed on the initial SDAY plates. This yeast isolate was grown subsequently at 28°C on pollen agar (1% pollen plus 1.5% agar), Lee's agar, Czapek-Dox broth, M40Y agar (a high sucrose, osmotic-stress medium), and moistened sterilized bee-collected pollen.

Yeast Fatty Acid Methyl Esters (FAMES). Cells of the SHB larval yeast isolate were harvested and treated chemically to extract and convert the fatty acids present in the cell wall or cell membrane fractions to FAMES (16). The total cellular FAMES were analyzed by GC and the resulting profiles matched with those of yeasts available in the Microbial Identification system

(MIDI) database by using Sherlock Version 4.5 software (Microbial ID, 1993) (17). The MIDI analysis identified the isolate as a close relative of *Candida krusei* producing a similarity index of 0.828 (16).

Yeast DNA Extraction, PCR, and Sequencing. The yeast isolate was inoculated in Sabouraud maltose broth and incubated overnight at 26°C. DNA was isolated from cell pellets by using the Masterpure yeast DNA purification kit (Epicentre, Madison, WI). The quantity and quality of the DNA was evaluated on ethidium-stained agarose gels. Aliquots of the DNA were amplified with a mixture of TaqDNA polymerase (Promega, Madison, WI) and *PFU* polymerase (Stratagene) by using the primers TW81 and AB28 for the ITS-5.8S (18) and NL-1 and NL-4 primers for the 28S (19). Both the D1/D2 and ITS1–5.8S-ITS2 sequences, when examined by BLAST analysis and database searches, produced matches with extremely low expect values. The 509-bp D1/D2 sequence (GenBank accession no. AY911384) was 100% homologous to four strains (accession nos. AF335976, AY267821, U45702, and AY267824) of *K. (Pichia) ohmeri*, an unidentified yeast species (AF335975), and to *Candida membranifaciens* (AJ508563). The ITS1–5.8S-ITS2 sequence (GenBank accession no. AY911385) was 99–100% homologous to various *K. ohmeri* strains (accession nos. AY168786, AF219004, and AF218977) and the unknown yeast isolates AF536211 and AF536209. The molecular data identified the fungal isolate as a strain of *K. ohmeri*.

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